IAP20 Rec'd PCT/PTO 25 MAY 2006

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<110>	Ono, Yuichi Nakagawa, Yasuko Nakatani, Tomoya	
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Glu Lys Ile Ala Pro Thr Glu Phe Val Met Arg Ala Leu Glu Cys Val
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Tyr His Leu Gly Cys Phe Cys Cys Cys Val Cys Glu Arg Gln Leu Arg
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Asp Tyr Glu Lys Glu Lys Asp Leu Leu Ser Ser Val Ser Pro Asp Glu
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Ser Asp Ser Val Lys Ser Glu Asp Glu Asp Gly Asp Met Lys Pro Ala
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Tyr Phe Ala Ser 370

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Glu Val Leu Ser Ser Arg Met Glu Gly Met Met Ala Ser Tyr Thr Pro
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Gly Ser Ser Asp Pro Phe Gln Gln Gly Leu Thr Pro Pro Gln Met Pro
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Gly Asn Asp Ser Ile Phe His Asp Ile Asp Ser Asp Thr Ser Leu Thr
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Asp Pro Gly Ser Leu His Asn Phe His Gln Asn Tyr Val Ala Thr Thr
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Glu Ser Gln Ile Lys Ile Trp Phe Gln Asn Lys Arg Ala Lys Ile Lys
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Lys Ala Thr Gly Ile Lys Asn Gly Leu Ala Leu His Leu Met Ala Gln
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Phe Ile Gly Arg Arg Gln Ser Leu Ile Glu Asp Ala Arg Lys Glu Arg
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Val Ile Leu Ile Ser Phe Tyr Val Gly Phe Phe Tyr Asn Val Ile Ile
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Trp Ile His Cys Asn Asn Thr Trp Asn Ser Pro Asn Cys Ser Asp Ala
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Thr Thr Pro Ala Ala Glu Tyr Phe Glu Arg Gly Val Leu His Leu His
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Tyr Met Ala Gln Lys His Asn Val Pro Ile Arg Asp Val Ala Thr Asp
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Thr Ser Ile Leu Phe Gly Val Leu Ile Glu Ala Ile Gly Val Ala Trp
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Tyr Lys Phe Cys Ser Leu Pro Gly Ser Phe Arg Glu Lys Leu Ala Tyr
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Lys Lys Ile Asp Phe Leu Leu Ser Val Ile Gly Phe Ala Val Asp Leu
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Ala Phe Leu Val Pro Tyr Leu Leu Phe Met Val Ile Ala Gly Met Pro
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Ala Gly Val Trp Lys Ile Cys Pro Ile Leu Lys Gly Val Gly Phe Thr
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28

140

155

170

135

150

165

180

195

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Ala Phe Lys Ala Ser Phe Glu Val Ser Ser Lys Pro Cys Arg Lys Val 210 215 220

Arg Glu Thr Leu Ala Ala Glu Thr Gly Leu Ser Val Arg Val Val Gin 225 230 235 240

Val Trp Phe Gin Asn Gin Arg Ala Lys Met Lys Lys Leu Ala Arg Arg
245 250 255

Gin Gin Gin Gin Gin Asp Gin Gin Asp Thr Gin Arg Leu Thr Ser 260 265 270

Ala Gin Thr Asn Gly Ser Gly Asn Ala Gly Met Glu Gly He Met Asn 275 280 285

Pro Tyr Thr Thr Leu Pro Thr Pro Gin Gin Leu Leu Ala IIe Giu Gin 290 295 300

Ser Val Tyr Asn Ser Asp Pro Phe Arg Gln Gly Leu	Thr Pro Pro GIn
305 310 315	320
Met Pro Gly Asp His Met His Pro Tyr Gly Ala Glu F	Pro Leu Phe His
325 330	335
·	
Asn Lau Asn Sar Asn Asn Thr Sar Lau Sar Asn Lau	No. Ann. Over Die
Asp Leu Asp Ser Asp Asp Thr Ser Leu Ser Asn Leu 6 340 345	350
3.0	
Leu Ala Thr Ser Glu Ala Gly Pro Leu Gln Ser Arg V	/al Gly Asn Pro
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300

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<211> 382

<212> PRT

<213> Homo sapiens

<400> 16

Met Leu Asp Gly Leu Lys Met Glu Glu Asn Phe Gln Ser Ala IIe Asp 1 5 10 15

Thr Ser Ala Ser Phe Ser Ser Leu Leu Gly Arg Ala Val Ser Pro Lys
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Ser Val Cys Glu Gly Cys Gln Arg Val IIe Leu Asp Arg Phe Leu Leu 35 40 45

Arg Leu Asn Asp Ser Phe Trp His Glu Gln Cys Val Gln Cys Ala Ser 50 55 60

Cys Lys Glu Pro Leu Glu Thr Thr Cys Phe Tyr Arg Asp Lys Lys Leu 65 70 75 80

Tyr Cys Lys Tyr Asp Tyr Glu Lys Leu Phe Ala Val Lys Cys Gly Gly
85 90 95

Cys Phe Glu Ala IIe Ala Pro Asn Glu Phe Val Met Arg Ala Gln Lys
100 105 110

Ser Val Tyr His Leu Ser Cys Phe Cys Cys Cys Val Cys Glu Arg Gln 115 120 125

Leu Gin Lys Gly Asp Glu Phe Val Leu Lys Glu Gly Gin Leu Leu Cys 130 135 140

Lys Gly Asp Tyr Glu Lys Glu Arg Glu Leu Leu Ser Leu Val Ser Pro 145 150 155 160

Ala Ala Ser Asp Ser Gly Lys Ser Asp Glu Glu Ser Leu Cys Lys
165 170 175

Ser Ala His Gly Ala Gly Lys Gly Thr Ala Glu Glu Gly Lys Asp His 180 185 190

Lys Arg Pro Lys Arg Pro Arg Thr IIe Leu Thr Thr Gln Gln Arg Arg
195 200 205

Ala Phe Lys Ala Ser Phe Glu Val Ser Ser Lys Pro Cys Arg Lys Val 210 215 220

Arg 225	Glu	Ihr	Leu	Ala	A1a 230	Glu	Thr	Gly	Leu	Ser 235	Val	Arg	Val	Val	GIn 240
										200					240
Val	Trp	Phe	Gln		GIn	Arg	Ala	Lys		Lys	Lys	Leu	Ala		Arg
				245					250					255	
GIn	GIn	Gin	GIn	GIn	GIn	Asp	GIn	Gin	Asn	Thr	Gln	Arg	Leu	Ser	Ser
			260					265					270		
Ala	Gin	Thr	Asn	Gly	Gly	Gly	Ser	Ala	Gly	Met	Glu	Gly	Пe	Met	Asn
		275					280					285			
Pro	Tyr	Thr	Ala	Leu	Pro	Thr	Pro	Gin	Gln	Leu	Leu	Ala	lle	Glu	Gln
	290					295					300				
Ser	Val	Tyr	Ser	Ser	Asp	Pro	Phe	Arg	Gln	Gly	Leu	Thr	Pro	Pro	Gln
305					310					315					320

Met Pro Gly Asp His Met His Pro Tyr Gly Ala Glu Pro Leu Phe His

330

335

325

Asp Leu Asp Ser Asp Asp Thr Ser Leu Ser Asn Leu Gly Asp Cys Phe 340 345 350

Leu Ala Thr Ser Glu Ala Gly Pro Leu Gln Ser Arg Val Gly Asn Pro 355 360 365 Ile Asp His Leu Tyr Ser Met Gln Asn Ser Tyr Phe Thr Ser 370 375 380

<210> 17

<211> 839

<212> DNA

<213≻ Homo sapiens

<400> 17

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taatgatatg gtgtagctca gcatttccaa agactgaata cattatggat tgcatagtt

839

<210> 18

<211> 133

<212> PRT

<213> Homo sapiens

<400> 18

Met Lys Lys Leu Ala Arg Arg Gin Gin Gin Gin Gin Gin Asp Gin Gin

1 15

Asn Thr Gln Arg Leu Ser Ser Ala Gln Thr Asn Gly Gly Ser Ala
20 25 30

Gly Met Glu Gly lle Met Asn Pro Tyr Thr Ala Leu Pro Thr Pro Gln 35 40 45

GIn Leu Leu Ala IIe Glu GIn Ser Val Tyr Ser Ser Asp Pro Phe Arg 50 55 60

Gln Gly Leu Thr Pro Pro Gln Met Pro Gly Asp His Met His Pro Tyr 65 70 75 80

Gly Ala Glu Pro Leu Phe His Asp Leu Asp Ser Asp Asp Thr Ser Leu 85 90 95

Ser Asn Leu Gly Asp Cys Phe Leu Ala Thr Ser Glu Ala Gly Pro Leu

100 105 110

Gln Ser Arg Val Gly Asn Pro IIe Asp His Leu Tyr Ser Met Gln Asn 115 120 125

Ser Tyr Phe Thr Ser 130

<210> 19

<211> 1119

<212> DNA

<213> Mus musculus

<400> 19

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accatcctca	ccacacagca	gcgaagagct	ttcaaggcat	cctttgaggt	ctcctccaag	660
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caggtctggt	ttcagaacca	aagagcaaag	atgaagaagc	tggcccggag	acaccagcaa	780
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ggcatgatgg	cctcctacac	cgcgctggcc	cctccgcagc	agcagatcgt	ggccatggag	900
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gggaacgact	ccatcttcca	cgatattgat	agtgatacct	ccctcaccag	cctcagcgac	1020
tgcttcctcg	gctcttccga	cgtgggctcc	ctgcaggcgc	gcgtggggaa	ccccattgac	1080
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<211> 372

<212> PRT

<213> Mus musculus

<400> 20

Met Leu Asp Gly Ile Lys Met Glu Glu His Ala Leu Arg Pro Gly Pro 1 5 10 15

Ala Thr Leu Gly Val Leu Leu Gly Ser Asp Cys Pro His Pro Ala Val 20 25 30

Cys Glu Gly Cys Gln Arg Pro IIe Ser Asp Arg Phe Leu Met Arg Val

35 40 45

Asn Glu Ser Ser Trp His Glu Glu Cys Leu Gln Cys Ala Ala Cys Gln
50 55 60

Gin Ala Leu Thr Thr Ser Cys Tyr Phe Arg Asp Arg Lys Leu Tyr Cys 65 70 75 80

Lys Gln Asp Tyr Gln Gln Leu Phe Ala Ala Lys Cys Ser Gly Cys Met 85 90 95

Glu Lys lie Ala Pro Thr Glu Phe Val Met Arg Ala Leu Glu Cys Val 100 105 110

Tyr His Leu Gly Cys Phe Cys Cys Cys Val Cys Glu Arg Gln Leu Arg 115 120 125

Lys Gly Asp Glu Phe Val Leu Lys Glu Gly Gln Leu Leu Cys Lys Gly
130 135 140

Asp Tyr Glu Lys Glu Lys Asp Leu Leu Ser Ser Val Ser Pro Asp Glu 145 150 155 160

Ser Asp Ser Val Lys Ser Glu Asp Glu Asp Gly Asp Met Lys Pro Ala 165 170 175

Lys Gly Gln Gly Ser Gln Ser Lys Gly Ser Gly Asp Asp Gly Lys Asp

180 185 190

Pro Arg Arg Pro Lys Arg Pro Arg Thr IIe Leu Thr Thr Gin Gin Arg
195 200 205

Arg Ala Phe Lys Ala Ser Phe Glu Val Ser Ser Lys Pro Cys Arg Lys 210 215 220

Val Arg Glu Thr Leu Ala Ala Glu Thr Gly Leu Ser Val Arg Val Val 225 230 235 240

Gin Val Trp Phe Gin Asn Gin Arg Ala Lys Met Lys Lys Leu Ala Arg 245 250 255

Arg His Gln Gln Gln Gln Gln Gln Asn Ser Gln Arg Leu Gly Gln 260 265 270

Glu Val Leu Ser Ser Arg Met Glu Gly Met Met Ala Ser Tyr Thr Ala 275 280 285

Leu Ala Pro Pro Gin Gin Gin lie Val Ala Met Giu Gin Ser Pro Tyr 290 295 300

Gly Ser Ser Asp Pro Phe Gln Gln Gly Leu Thr Pro Pro Gln Met Pro 305 310 315 320

Gly Asn Asp Ser lie Phe His Asp Ile Asp Ser Asp Thr Ser Leu Thr

325 330 335

Ser Leu Ser Asp Cys Phe Leu Gly Ser Ser Asp Val Gly Ser Leu Gln 340 345 350

Ala Arg Val Gly Asn Pro lle Asp Arg Leu Tyr Ser Met Gln Ser Ser 355 360 365

Tyr Phe Ala Ser 370

<210> 21

<211> 1119

<212> DNA

<213> Homo sapiens

<400> 21

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ccttgccgaa	aggtccgaga	gacactggca	gctgagacgg	gcctcagtgt	gcgcgtggtc	720
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cagagcccct	acggcagcag	cgaccccttc	cagcagggcc	tcacgccgcc	ccaaatgcca	960
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tgcttcctcg	gctcctcaga	cgtgggctcc	ctgcaggccc	gcgtggggaa	ccccatcgac	1080
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⟨211⟩ 372

<212> PRT

<213> Homo sapiens

<400> 22

Met Leu Asp Gly lle Lys Met Glu Glu His Ala Leu Arg Pro Gly Pro 1 5 10 15

Ala Thr Leu Gly Val Leu Leu Gly Ser Asp Cys Pro His Pro Ala Val

20 25 30

Cys Glu Gly Cys Gln Arg Pro lle Ser Asp Arg Phe Leu Met Arg Val 35 40 45

Asn Glu Ser Ser Trp His Glu Glu Cys Leu Gln Cys Ala Ala Cys Gln 50 55 60

Gin Ala Leu Thr Thr Ser Cys Tyr Phe Arg Asp Arg Lys Leu Tyr Cys
65 70 75 80

Lys Gin Asp Tyr Gin Gin Leu Phe Ala Ala Lys Cys Ser Giy Cys Met 85 90 95

Glu Lys IIe Ala Pro Thr Glu Phe Val Met Arg Ala Leu Glu Cys Val 100 105 110

Tyr His Leu Gly Cys Phe Cys Cys Cys Val Cys Glu Arg Gln Leu Arg
115 120 125

Lys Gly Asp Glu Phe Val Leu Lys Glu Gly Gln Leu Leu Cys Lys Gly
130 135 140

Asp Tyr Glu Lys Glu Lys Asp Leu Leu Ser Ser Val Ser Pro Asp Glu 145 150 155 160

Ser Asp Ser Val Lys Ser Glu Asp Glu Asp Gly Asp Met Lys Pro Ala

165 170 175

Lys Gly Gln Gly Ser Gln Ser Lys Gly Ser Gly Asp Asp Gly Lys Asp 180 185 190

Pro Arg Arg Pro Lys Arg Pro Arg Thr IIe Leu Thr Thr Gin Gin Arg 195 200 205

Arg Ala Phe Lys Ala Ser Phe Glu Val Ser Ser Lys Pro Cys Arg Lys
210 215 220

Val Arg Glu Thr Leu Ala Ala Glu Thr Gly Leu Ser Val Arg Val Val 225 230 235 240

Gin Val Trp Phe Gin Asn Gin Arg Ala Lys Met Lys Lys Leu Ala Arg 245 250 255

Arg His Gln Gln Gln Gln Gln Gln Asn Ser Gln Arg Leu Gly Gln
260 265 270

Glu Val Leu Ser Ser Arg Met Glu Gly Met Met Ala Ser Tyr Thr Pro 275 280 285

Leu Ala Pro Pro Gin Gin Gin lie Val Ala Met Giu Gin Ser Pro Tyr 290 295 300

Gly Ser Ser Asp Pro Phe Gln Gln Gly Leu Thr Pro Pro Gln Met Pro

305 310 315 320

Gly Asn Asp Ser IIe Phe His Asp IIe Asp Ser Asp Thr Ser Leu Thr
325 330 335

Ser Leu Ser Asp Cys Phe Leu Gly Ser Ser Asp Val Gly Ser Leu Gln 340 345 350

Ala Arg Val Gly Asn Pro IIe Asp Arg Leu Tyr Ser Met Gln Ser Ser 355 360 365

Tyr Phe Ala Ser 370

<210> 23

<211> 2247

<212> DNA

<213> Mus musculus

<400> 23

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ccgcttccgc agcgctcccg cggacccggg ctcctctgct cccggaggga actgcacttc 180
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aatttcctcg aaaactccaa taactctgct gaagccatgc cttgtgtca ggcgcagtat 360

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actgaaatta	ctgccaccac	ttctctcccc	agcttcagta	cctttatgga	caactacagc	540
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gacaaacttt	tcctggacac	cttacctttc	taagaccttc	tcccaagcac	gtcaaagaac	2160
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aagtgtccag	ttcaccttat	ctccctt				2247

<211> 598

<212> PRT

<213> Mus musculus

<400> 24

Met Pro Cys Val Gln Ala Gln Tyr Gly Ser Ser Pro Gln Gly Ala Ser

1 5 10 15

Pro Ala Ser Gln Ser Tyr Ser Tyr His Ser Ser Gly Glu Tyr Ser Ser 20 25 30

Asp Phe Leu Thr Pro Glu Phe Val Lys Phe Ser Met Asp Leu Thr Asn 35 40 45

Thr Glu IIe Thr Ala Thr Thr Ser Leu Pro Ser Phe Ser Thr Phe Met 50 55 60

Asp Asn Tyr Ser Thr Gly Tyr Asp Val Lys Pro Pro Cys Leu Tyr Gln 65 70 75 80

Met Pro Leu Ser Gly Gln Gln Ser Ser Ile Lys Val Glu Asp Ile Gln 85 90 95

Met His Asn Tyr Gln Gln His Ser His Leu Pro Pro Gln Ser Glu Glu
100 105 110

Met Met Pro His Ser Gly Ser Val Tyr Tyr Lys Pro Ser Ser Pro Pro 115 120 125

Thr Pro Ser Thr Pro Ser Phe Gin Val Gin His Ser Pro Met Trp Asp 130 135 140 Asp Pro Gly Ser Leu His Asn Phe His Gln Asn Tyr Val Ala Thr Thr 145 150 155 160

His Met IIe Glu Gln Arg Lys Thr Pro Val Ser Arg Leu Ser Leu Phe 165 170 175

Ser Phe Lys Gln Ser Pro Pro Gly Thr Pro Val Ser Ser Cys Gln Met 180 185 190

Arg Phe Asp Gly Pro Leu His Val Pro Met Asn Pro Glu Pro Ala Gly
195 200 205

Ser His His Val Val Asp Gly Gln Thr Phe Ala Val Pro Asn Pro IIe 210 215 220

Arg Lys Pro Ala Ser Met Gly Phe Pro Gly Leu Gln IIe Gly His Ala 225 230 235 240

Ser Gln Leu Leu Asp Thr Gln Val Pro Ser Pro Pro Ser Arg Gly Ser

245 250 255

Pro Ser Asn Glu Gly Leu Cys Ala Val Cys Gly Asp Asn Ala Ala Cys 260 265 270

Gln His Tyr Gly Val Arg Thr Cys Glu Gly Cys Lys Gly Phe Phe Lys 275 280 285 Arg Thr Val Gln Lys Asn Ala Lys Tyr Val Cys Leu Ala Asn Lys Asn 290 295 300

Cys Pro Val Asp Lys Arg Arg Arg Asn Arg Cys Gin Tyr Cys Arg Phe 305 310 315 320

Gin Lys Cys Leu Ala Val Gly Met Val Lys Glu Val Val Arg Thr Asp 325 330 335

Ser Leu Lys Gly Arg Arg Gly Arg Leu Pro Ser Lys Pro Lys Ser Pro 340 345 350

Gln Asp Pro Ser Pro Pro Ser Pro Pro Val Ser Leu IIe Ser Ala Leu 355 360 365

Val Arg Ala His Val Asp Ser Asn Pro Ala Met Thr Ser Leu Asp Tyr 370 375 380

Ser Arg Phe Gln Ala Asn Pro Asp Tyr Gln Met Ser Gly Asp Asp Thr 385 390 395 400

Gln His IIe Gln Gln Phe Tyr Asp Leu Leu Thr Gly Ser Met Glu IIe
405 410 415

Ile Arg Gly Trp Ala Glu Lys Ile Pro Gly Phe Ala Asp Leu Pro Lys
420 425 430

Ala Asp Gln Asp Leu Leu Phe Glu Ser Ala Phe Leu Glu Leu Phe Val 435 440 445

Leu Arg Leu Ala Tyr Arg Ser Asn Pro Vai Glu Gly Lys Leu lle Phe 450 455 460

Cys Asn Gly Val Val Leu His Arg Leu Gln Cys Val Arg Gly Phe Gly 465 470 475 480

Glu Trp IIe Asp Ser IIe Val Glu Phe Ser Ser Asn Leu Gln Asn Met 485 490 495

Asn Ile Asp Ile Ser Ala Phe Ser Cys Ile Ala Ala Leu Ala Met Val 500 505 510

Thr Glu Arg His Gly Leu Lys Glu Pro Lys Arg Val Glu Glu Leu Gln
515 520 525

Asn Lys lle Val Asn Cys Leu Lys Asp His Val Thr Phe Asn Asn Gly 530 535 540

Gly Leu Asn Arg Pro Asn Tyr Leu Ser Lys Leu Leu Gly Lys Leu Pro 545 550 555 560

Glu Leu Arg Thr Leu Cys Thr Gln Gly Leu Gln Arg IIe Phe Tyr Leu
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Leu Asp Thr Leu Pro Phe 595

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<213> Homo sapiens

<400> 25

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<213> Homo sapiens

<400> 26

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Asp Phe Leu Thr Pro Glu Phe Val Lys Phe Ser Met Asp Leu Thr Asn 35 40 45

Thr Glu lle Thr Ala Thr Thr Ser Leu Pro Ser Phe Ser Thr Phe Met 50 55 60

Asp Asn Tyr Ser Thr Gly Tyr Asp Val Lys Pro Pro Cys Leu Tyr Gln 65 70 75 80

Met Pro Leu Ser Gly Gln Gln Ser Ser Ile Lys Val Glu Asp Ile Gln 85 90 95

Met His Asn Tyr Gln Gln His Ser His Leu Pro Pro Gln Ser Glu Glu
100 105 110

Met Met Pro His Ser Gly Ser Val Tyr Tyr Lys Pro Ser Ser Pro Pro 115 120 125

Thr Pro Thr Thr Pro Gly Phe Gln Val Gln His Ser Pro Met Trp Asp 130 135 140

Asp Pro Gly Ser Leu His Asn Phe His Gln Asn Tyr Val Ala Thr Thr 145 150 155 160

His Met Ile Glu Gln Arg Lys Thr Pro Val Ser Arg Leu Ser Leu Phe

165 170 175

Ser Phe Lys Gin Ser Pro Pro Gly Thr Pro Val Ser Ser Cys Gin Met 180 185 190

Arg Phe Asp Gly Pro Leu His Val Pro Met Asn Pro Glu Pro Ala Gly
195 200 205

Ser His His Val Val Asp Gly Gln Thr Phe Ala Val Pro Asn Pro IIe 210 215 220

Arg Lys Pro Ala Ser Met Gly Phe Pro Gly Leu Gln Ile Gly His Ala 225 230 235 240

Ser Gin Leu Leu Asp Thr Gin Val Pro Ser Pro Pro Ser Arg Gly Ser 245 250 255

Pro Ser Asn Glu Gly Leu Cys Ala Val Cys Gly Asp Asn Ala Ala Cys 260 265 270

Gln His Tyr Gly Val Arg Thr Cys Glu Gly Cys Lys Gly Phe Phe Lys 275 280 285

Arg Thr Val Gin Lys Asn Ala Lys Tyr Val Cys Leu Ala Asn Lys Asn 290 295 300

Cys Pro Val Asp Lys Arg Arg Arg Asn Arg Cys Gln Tyr Cys Arg Phe

305 310 315 320

Gin Lys Cys Leu Ala Val Gly Met Val Lys Glu Val Val Arg Thr Asp 325 330 335

Ser Leu Lys Gly Arg Arg Gly Arg Leu Pro Ser Lys Pro Lys Ser Pro 340 345 350

Gin Giu Pro Ser Pro Pro Ser Pro Pro Val Ser Leu IIe Ser Ala Leu 355 360 365

Val Arg Ala His Val Asp Ser Asn Pro Ala Met Thr Ser Leu Asp Tyr 370 375 380

Ser Arg Phe Gln Ala Asn Pro Asp Tyr Gln Met Ser Gly Asp Asp Thr 385 390 395 400

GIn His IIe GIn GIn Phe Tyr Asp Leu Leu Thr Gly Ser Met Glu IIe 405 410 415

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420 425 430

Ala Asp Gin Asp Leu Leu Phe Giu Ser Ala Phe Leu Giu Leu Phe Val 435 440 445

Leu Arg Leu Ala Tyr Arg Ser Asn Pro Val Glu Gly Lys Leu lle Phe

450 455 460

Cys Asn Gly Val Val Leu His Arg Leu Gln Cys Val Arg Gly Phe Gly 465 470 475 480

Glu Trp lle Asp Ser lle Val Glu Phe Ser Ser Asn Leu Gln Asn Met
485 490 495

Asn IIe Asp IIe Ser Ala Phe Ser Cys IIe Ala Ala Leu Ala Met Val 500 505 510

Thr Glu Arg His Gly Leu Lys Glu Pro Lys Arg Val Glu Glu Leu Gln
515 520 525

Asn Lys IIe Val Asn Cys Leu Lys Asp His Val Thr Phe Asn Asn Giy 530 535 540

Gly Leu Asn Arg Pro Asn Tyr Leu Ser Lys Leu Leu Gly Lys Leu Pro 545 550 555 560

Glu Leu Arg Thr Leu Cys Thr Gln Gly Leu Gln Arg lle Phe Tyr Leu
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Lys Leu Glu Asp Leu Vai Pro Pro Pro Ala IIe IIe Asp Lys Leu Phe 580 585 590 595

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<400> 28

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Pro Gly Ala Ser Gly Ser Ser Gly Ser Asp Gly Asp Ser Val Pro Val
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Ser Pro Gln Pro Ala Pro Pro Ser Pro Pro Ala Ala Pro Cys Leu Pro Pro Leu Ala His His Pro His Leu Pro Pro His Pro Gln His Leu Ala Ala Pro Ala His Gln Pro Gln Pro Ala Ala Gln Leu His Arg Thr Thr Asn Phe Phe Ile Asp Asn Ile Leu Arg Pro Asp Phe Gly Cys Lys Lys Glu Gin Pro Leu Pro Gin Leu Leu Val Ala Ser Ala Ala Ala Gly Gly Gly Ala Ala Ala Gly Gly Gly Ser Arg Val Glu Arg Asp Arg Gly Gln Thr Gly Ala Gly Arg Asp Pro Val His Ser Leu Gly Thr Arg Ala Ser Gly Ala Ala Ser Leu

Leu Cys Ala Pro Asp Ala Asn Cys Gly Pro Pro Asp Gly Ser Gln Pro
180 185 190

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Arg	Tyr	Ser	Asp	Arg	Pro	Ser	Ser	Gly	Pro	Arg	Thr	Arg	Lys	Leu	Lys
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Lys Lys Lys Asn Glu Lys Glu Asp Lys Arg Pro Arg Thr Ala Phe Thr

315

320

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Ile Thr Glu Gln Arg Arg Gln Thr Leu Ala Gln Glu Leu Ser Leu Asn 340 345 350

Glu Ser Gln lie Lys lie Trp Phe Gln Asn Lys Arg Ala Lys lie Lys 355 360 365

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<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

<400> 30

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Pro Gly Ala Ser Gly Ser Ser Gly Ser Gly Ser Asp Gly Asp Ser Val

35 40 45

Pro Val Ser Pro Gin Pro Ala Pro Pro Ser Pro Pro Ala Ala Pro Cys
50 55 60

Leu Pro Pro Leu Ala His His Pro His Leu Pro Pro His Pro Pro Pro 65 70 75 80

Pro Pro Pro Gln His Leu Ala Ala Pro Ala His Gln Pro Gln Pro Ala 85 90 95

Ala Gin Leu His Arg Thr Thr Asn Phe Phe IIe Asp Asn IIe Leu Arg 100 105 110

Pro Asp Phe Gly Cys Lys Lys Glu Gln Pro Pro Pro Gln Leu Leu Val 115 120 125

Ala Ala Ala Arg Gly Gly Ala Gly Gly Gly Gly Arg Val Glu Arg 130 135 140

Asp Arg Gly Gln Thr Ala Ala Gly Arg Asp Pro Val His Pro Leu Gly 145 150 155 160

Thr Arg Ala Pro Gly Ala Ala Ser Leu Leu Cys Ala Pro Asp Ala Asn 165 170 175

Cys Gly Pro Pro Asp Gly Ser Gln Pro Ala Ala Ala Gly Ala Gly Ala

180 185 190

Val Ala Ala Ala Ala Ala Ala Ala Ala Lys Pro Ser Asp Thr Gly 210 215 220

Gly Gly Gly Ser Gly Gly Gly Ala Gly Ser Pro Gly Ala Gln Gly Thr 225 230 235 240

Lys Tyr Pro Glu His Gly Asn Pro Ala IIe Leu Leu Met Gly Ser Ala 245 250 255

Asn Gly Gly Pro Val Val Lys? Thr Asp Ser Gln Gln Pro Leu Val Trp
260 265 270

Pro Ala Trp Val Tyr Cys Thr Arg Tyr Ser Asp Arg Pro Ser Ser Gly
275 280 285

Pro Arg Thr Arg Lys Leu Lys Lys Lys Asn Glu Lys Glu Asp Lys 290 295 300

Arg Pro Arg Thr Ala Phe Thr Ala Glu Gln Leu Gln Arg Leu Lys Ala 305 310 315 320

Glu Phe Gln Ala Asn Arg Tyr lle Thr Glu Gln Arg Arg Gln Thr Leu

325 330 335

Ala Gin Giu Leu Ser Leu Asn Giu Ser Gin IIe Lys IIe Trp Phe Gin 340 345 350

Asn Lys Arg Ala Lys IIe Lys Lys Ala Thr Gly IIe Lys Asn Gly Leu 355 360 365

Ala Leu His Leu Met Ala Gln Gly Leu Tyr Asn His Ser Thr Thr Thr 370 375 380

Val Gln Asp Lys Asp Glu Ser Glu 385 390

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<212> DNA

<213> Mus musculus

<400> 31

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<210> 32

<211> 302

<212> PRT

<213> Mus musculus

<400> 32

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Ser Leu Ser Asp Ala Gly Thr Pro His Pro Pro Leu Pro Glu His Gly
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Cys Lys Gly Gln Glu His Ser Asp Ser Glu Lys Ala Ser Ala Ser Leu 35 40 45

Pro Gly Gly Ser Pro Glu Asp Gly Ser Leu Lys Lys Gln Arg Arg
50 55 60

GIn Arg Thr His Phe Thr Ser GIn GIn Leu GIn Glu Leu GIu Ala Thr 65 70 75 80

Phe Gin Arg Asn Arg Tyr Pro Asp Met Ser Thr Arg Giu Giu ile Ala 85 90 95

Val Trp Thr Asn Leu Thr Glu Ala Arg Val Arg Val Trp Phe Lys Asn 100 105 110 Arg Arg Ala Lys Trp Arg Lys Arg Glu Arg Ser Gln Gln Ala Glu Leu 115 120 125

Cys Lys Gly Gly Phe Ala Ala Pro Leu Gly Gly Leu Val Pro Pro Tyr 130 135 140

Glu Glu Val Tyr Pro Gly Tyr Ser Tyr Gly Asn Trp Pro Pro Lys Ala 145 150 155 160

Leu Ala Pro Pro Leu Ala Ala Lys Thr Phe Pro Phe Ala Phe Asn Ser 165 170 175

Val Asn Val Gly Pro Leu Ala Ser Gln Pro Val Phe Ser Pro Pro Ser 180 185 190

Ser IIe Ala Ala Ser Met Val Pro Ser Ala Ala Ala Ala Pro Gly Thr 195 200 205

Val Pro Gly Pro Gly Ala Leu Gln Gly Leu Gly Gly Ala Pro Pro Gly 210 215 220

Leu Ala Pro Ala Ala Val Ser Ser Gly Ala Val Ser Cys Pro Tyr Ala 225 230 235 240

Ser Ala Ala Ala Ala Ala Ala Ala Ala Ser Ser Pro Tyr Val Tyr 245 250 255

Arg Asp Pro Cys Asn Ser Ser Leu Ala Ser Leu Arg Leu Lys Ala Lys
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Gin His Ala Ser Phe Ser Tyr Pro Ala Val Pro Gly Pro Pro Pro Ala 275 280 285

Ala Asn Leu Ser Pro Cys Gin Tyr Ala Vai Giu Arg Pro Vai 290 295 300

<210> 33

<211> 1407

<212> DNA

<213> Homo sapiens

<400> 33

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⟨210⟩ 34

⟨211⟩ 302

<212> PRT

<213≻ Homo sapiens

<400> 34

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Cys Lys Gly Gln Glu His Ser Asp Ser Glu Lys Ala Ser Ala Ser Leu 35 40 45

Pro Gly Gly Ser Pro Glu Asp Gly Ser Leu Lys Lys Lys Gln Arg Arg 50 55 60

GIn Arg Thr His Phe Thr Ser GIn GIn Leu GIn Glu Leu GIu Ala Thr 65 70 75 80

Phe Gln Arg Asn Arg Tyr Pro Asp Met Ser Thr Arg Glu Glu IIe Ala 85 90 95

Val Trp Thr Asn Leu Thr Glu Ala Arg Val Arg Val Trp Phe Lys Asn 100 105 110

Arg Arg Ala Lys Trp Arg Lys Arg Glu Arg Ser Gln Gln Ala Glu Leu 115 120 125

Cys Lys Gly Ser Phe Ala Ala Pro Leu Gly Gly Leu Val Pro Pro Tyr

130 135 140

Glu Glu Val Tyr Pro Gly Tyr Ser Tyr Gly Asn Trp Pro Pro Lys Ala

145 150 155 160

Leu Ala Pro Pro Leu Ala Ala Lys Thr Phe Pro Phe Ala Phe Asn Ser 165 170 175

Val Asn Val Gly Pro Leu Ala Ser Gln Pro Val Phe Ser Pro Pro Ser 180 185 190

Ser IIe Ala Ala Ser Met Val Pro Ser Ala Ala Ala Ala Pro Gly Thr
195 200 205

Val Pro Gly Pro Gly Ala Leu Gln Gly Leu Gly Gly Gly Pro Pro Gly
210 215 220

Leu Ala Pro Ala Ala Val Ser Ser Gly Ala Val Ser Cys Pro Tyr Ala 225 230 235 240

Ser Ala Ala Ala Ala Ala Ala Ala Ala Ser Ser Pro Tyr Val Tyr
245 250 255

Arg Asp Pro Cys Asn Ser Ser Leu Ala Ser Leu Arg Leu Lys Ala Lys
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GIn His Ala Ser Phe Ser Tyr Pro Ala Val His Gly Pro Pro Pro Ala

275 280 285

Ala Asn Leu Ser Pro Cys Gln Tyr Ala Val Glu Arg Pro Val 290 295 300

<210> 35

<211> 1757

<212> DNA

<213> Mus musculus

<400> 35

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Leu Ser Arg Ala Leu Lys Val Phe Glu Thr Phe Glu Ala Lys Ile His 100 105 110

90

Hj s	Leu	Glu	Thr	Arg	Pro	Ala	Gin	Arg	Pro	Leu	Ala	Gly	Ser	Pro	His
		115					120					125			

Leu	Glu	Tyr	Phe	Val	Arg	Phe	Glu	Val	Pro	Ser	Gly	Asp	Leu	Ala	Ala
	130					135					140				

Leu Leu Ser Ser Val Arg Arg Val Ser Asp Asp Val Arg Ser Ala Arg 145 150 155 160

Glu Asp Lys Val Pro Trp Phe Pro Arg Lys Val Ser Glu Leu Asp Lys
165 170 175

Cys His His Leu Val Thr Lys Phe Asp Pro Asp Leu Asp Leu Asp His 180 185 190

Pro Gly Phe Ser Asp Gln Ala Tyr Arg Gln Arg Arg Lys Leu IIe Ala 195 200 205

Glu lle Ala Phe Gln Tyr Lys Gln Gly Glu Pro lle Pro His Val Glu 210 215 220

Tyr Thr Lys Glu Glu IIe Ala Thr Trp Lys Glu Val Tyr Ala Thr Leu 225 230 235 240

Lys Gly Leu Tyr Ala Thr His Ala Cys Arg Glu His Leu Glu Ala Phe 245 250 255 Gin Leu Leu Giu Arg Tyr Cys Giy Tyr Arg Giu Asp Ser ile Pro Gin 260 265 270

Leu Glu Asp Val Ser His Phe Leu Lys Glu Arg Thr Gly Phe Gln Leu 275 280 285

Arg Pro Val Ala Giy Leu Leu Ser Ala Arg Asp Phe Leu Ala Ser Leu 290 295 300

Ala Phe Arg Val Phe Gln Cys Thr Gln Tyr lle Arg His Ala Ser Ser 305 310 315 320

Pro Met His Ser Pro Glu Pro Asp Cys Cys His Glu Leu Leu Gly His
325 330 335

Val Pro Met Leu Ala Asp Arg Thr Phe Ala Gin Phe Ser Gin Asp Ile 340 345 350

Gly Leu Ala Ser Leu Gly Ala Ser Asp Glu Glu lle Glu Lys Leu Ser 355 360 365

Thr Val Tyr Trp Phe Thr Val Glu Phe Gly Leu Cys Lys Gln Asn Gly 370 375 380

Glu Leu Lys Ala Tyr Gly Ala Gly Leu Leu Ser Ser Tyr Gly Glu Leu 385 390 395 400 Leu His Ser Leu Ser Glu Glu Pro Glu Val Arg Ala Phe Asp Pro Asp 405 410 415

Thr Ala Ala Val Gln Pro Tyr Gln Asp Gln Thr Tyr Gln Pro Val Tyr 420 425 430

Phe Val Ser Glu Ser Phe Ser Asp Ala Lys Asp Lys Leu Arg Asn Tyr 435 440 445

Ala Ser Arg IIe Gin Arg Pro Phe Ser Val Lys Phe Asp Pro Tyr Thr 450 455 460

Leu Ala IIe Asp Val Leu Asp Ser Pro His Thr IIe Arg Arg Ser Leu 465 470 475 480

Glu Gly Val Gln Asp Glu Leu His Thr Leu Thr Gln Ala Leu Ser Ala 485 490 495

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<210> 37

<211> 1921

<212> DNA

<213> Homo sapiens

<400> 37

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<210> 38

<211> 528

<212> PRT

<213> Homo sapiens

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Arg Ala Val Lys Val Phe Glu Thr Phe Glu Ala Lys Ile His His Leu 130 135 140

Glu Thr Arg Pro Ala Gln Arg Pro Arg Ala Gly Gly Pro His Leu Glu 145 150 155 160

Tyr Phe Val Arg Leu Glu Val Arg Arg Gly Asp Leu Ala Ala Leu Leu
165 170 175

Ser Gly Val Arg Gln Val Ser Glu Asp Val Arg Ser Pro Ala Gly Pro 180 185 190

Lys Val Pro Trp Phe Pro Arg Lys Val Ser Glu Leu Asp Lys Cys His
195 200 205

His Leu Val Thr Lys Phe Asp Pro Asp Leu Asp Leu Asp His Pro Gly 210 215 220

Phe Ser Asp Gin Val Tyr Arg Gin Arg Arg Lys Leu IIe Ala Giu IIe 225 230 235 240

Ala Phe Gln Tyr Arg His Gly Asp Pro IIe Pro Arg Val Glu Tyr Thr
245 250 255

Ala Glu Glu IIe Ala Thr Trp Lys Glu Val Tyr Thr Thr Leu Lys Gly
260 265 270

Leu Tyr Ala Thr His Ala Cys Gly Glu His Leu Glu Ala Phe Ala Leu 275 280 285 Leu Glu Arg Phe Ser Gly Tyr Arg Glu Asp Asn Ile Pro Gln Leu Glu 290 295 300

Asp Val Ser Arg Phe Leu Lys Glu Arg Thr Gly Phe Gln Leu Arg Pro 305 310 315 320

Val Ala Gly Leu Leu Ser Ala Arg Asp Phe Leu Ala Ser Leu Ala Phe 325 330 335

Arg Val Phe Gln Cys Thr Gln Tyr lle Arg His Ala Ser Ser Pro Met 340 345 350

His Ser Pro Glu Pro Asp Cys Cys His Glu Leu Leu Gly His Val Pro 355 360 365

Met Leu Ala Asp Arg Thr Phe Ala Gin Phe Ser Gin Asp ile Gly Leu 370 375 380

Ala Ser Leu Gly Ala Ser Asp Glu Glu lle Glu Lys Leu Ser Thr Leu 385 390 395 400

Ser Trp Phe Thr Val Glu Phe Gly Leu Cys Lys Gln Asn Gly Glu Val
405 410 415

Lys Ala Tyr Gly Ala Gly Leu Leu Ser Ser Tyr Gly Glu Leu Leu His
420 425 430

Cys	Leu	Ser 435	Glu	Glu	Pro	Glu	lle 440	Arg	Ala	Phe	Asp	Pro 445	Glu	Ala	Ala	
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Ser 465	Glu	Ser	Phe	Ser	Asp 470	Ala	Lys	Asp	Lys	Leu 475	Arg	Ser	Tyr	Ala	Ser 480	
Arg	He	Gln	Arg	Pro 485	Phe	Ser	Val	Lys	Phe 490	Asp	Pro	Tyr	Thr	Leu 495	Ala	
He	Asp	Val	Leu 500	Asp	Ser	Pro	GIn	Ala 505	Val	Arg	Arg	Ser	Leu 510	Glu	Gly	
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120

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Ala Asn Val Trp Arg Phe Pro Tyr Leu Cys Tyr Lys Asn Gly Gly Gly

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Val lle Leu lle Ser Phe Tyr Val Gly Phe Phe Tyr Asn Val lle lle

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Ala Trp Ala Leu His Tyr Phe Phe Ser Ser Phe Thr Met Asp Leu Pro 165 170 175

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Cys Glu Ala Ser Val Trp I le Asp Ala Ala Thr Gln Val Cys Phe Ser 305 310 315 320

Leu Gly Val Gly Phe Gly Val Leu lle Ala Phe Ser Ser Tyr Asn Lys 325 330 335

Phe Thr Asn Asn Cys Tyr Arg Asp Ala lle lle Thr Thr Ser lle Asn 340 345 350

Ser Leu Thr Ser Phe Ser Ser Gly Phe Val Val Phe Ser Phe Leu Gly 355 360 365

Tyr Met Ala Gln Lys His Asn Val Pro IIe Arg Asp Val Ala Thr Asp 370 375 380

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Gin Arg Pro Asn Leu Tyr Trp Arg Leu Cys Trp Lys Leu Val Ser Pro 515 520 525

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Asn Pro Arg Gin Ser Pro Val Glu Ala Gin Asp Arg Glu Thr Trp Gly

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Lys Lys IIe Asp Phe Leu Leu Ser Val IIe Gly Phe Ala Val Asp Leu 65 70 75 80

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Ala Phe Leu Val Pro Tyr Leu Leu Phe Met Val IIe Ala Gly Met Pro 100 105 110

Leu Phe Tyr Met Glu Leu Ala Leu Gly Gln Phe Asn Arg Glu Gly Ala 115 120 125

Ala Gly Val Trp Lys lle Cys Pro lle Leu Lys Gly Val Gly Phe Thr 130 135 140

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His Gln Ser His Gly IIe Asp Asp Leu Gly Pro Pro Arg Trp Gln Leu 225 230 235 240

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Ser Leu Gly Val Gly Phe Gly Val Leu IIe Ala Phe Ser Ser Tyr Asn 325 330 335

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Thr Leu Phe IIe Val Leu Ala Thr Phe Leu Leu Ser Leu Phe Cys Val 450 455 460

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Gly Thr Ser lie Leu Phe Gly Val Leu lie Glu Ala lie Gly Val Ala

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Pro Cys Phe Leu Leu Phe Val Val Val Ser IIe Val Thr Phe Arg 530 535 540

Pro Pro His Tyr Gly Ala Tyr lle Phe Pro Asp Trp Ala Asn Ala Leu 545 550 555 560

Gly Trp Val lle Ala Thr Ser Ser Met Ala Met Val Pro lle Tyr Ala 565 570 575

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Leu Cys Gln Val Glu Glu Gly Asp Lys Glu Asp Val Asp Lys Ala Val
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Asp Ala Ser Glu Arg Gly Arg Leu Leu Tyr Lys Leu Ala Asp Leu IIe 85 90 95

Giu Arg Asp Arg Leu Leu Leu Ala Thr Met Giu Ser Met Asn Gly Gly
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Ala Ala Ser Arg IIe Phe Val Glu Glu Ser IIe Tyr Asp Glu Phe Val 305 310 315 320

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Glu Cys Gly Gly Gly Pro Trp Gly Asn Lys Gly Tyr Phe Val Gln Pro 370 375 380

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